RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
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form:

Application Serial Number: _10/612,468_

Source: _OIPE_

Date Processed by STIC: _7/29/2003_

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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
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MARK SPENCER, 703-308-4212.

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Revised 04/24/2003

OIPE

RAW SEQUENCE LISTING   DATE: 07/29/2003
PATENT APPLICATION: **US/10/612,468**   TIME: 13:35:30

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\07292003\J612468.raw**

```
        3 <110> APPLICANT: Zhang, Jingwu Z.
        4        Ho, Walter Kowk Keung
        5        Zhang, Dongqing
        6        Sun, Wei
W-->    7 <120> TITLE OF INVENTION: T Cell Receptor CDR3 Sequence and Methods for Detecting and
Treating
W-->    8        Rheumatoid Arthritis
W-->    9 <130> FILE REFERENCE: 057186.000003
W-->   10 <140> CURRENT APPLICATION NUMBER:
C-->   11 <141> CURRENT FILING DATE: 2003-07-02    ></1507
W-->   11 <151> PRIOR FILING DATE: 2002-07-02
W-->   12 <160> NUMBER OF SEQ ID: 168
```

← insert this mandatory numeric identifier and its response wherever </1517 for a response

Does Not Comply
Corrected Diskette Needed *pp 1,3*

ERRORED SEQUENCES

```
      775 <210> SEQ ID NO: 66    shown below
      776 <211> LENGTH: (21) 19
      777 <212> TYPE: DNA
      778 <213> ORGANISM: Artificial
W-->  779 <220> FEATURE:
      780 <221> NAME/KEY: primer_bind
      781 <223> OTHER INFORMATION: FAM (expand)-labeled BJ primer used for run-off reactions
W-->  782 <400> SEQUENCE: 66
E-->  784 ggagagtcga gttccatca                              19
     1416 <210> SEQ ID NO: 114
     1417 <211> LENGTH: 63
     1418 <212> TYPE: DNA
     1419 <213> ORGANISM: Artificial
W--> 1420 <220> FEATURE:
W--> 1421 <221> NAME/KEY:
     1422 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
     1423        RA patients
W--> 1424 <400> SEQUENCE: 114
     1426 tacttctgtg ccagcagtcc cacgcgggac aggggaaata atgagcagtt cttcgggcca    60
E--> 1427 gga                                                             63 ← insert
     1445 <210> SEQ ID NO: 116
     1446 <211> LENGTH: 63
     1447 <212> TYPE: DNA
     1448 <213> ORGANISM: Artificial    p.3
W--> 1449 <220> FEATURE:
W--> 1450 <221> NAME/KEY:
     1451 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
```

ST specimen of
     1452   RA patients

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W--> 1453 <400> SEQUENCE: 116
    1455 tacttctgtg ccagcagttc cccaatagcg gggagctcca atgagcagtt cttcgggcca   60 .
E--> 1456 gga   63 ← insert
    1473 <210> SEQ ID NO: 118
    1474 <211> LENGTH: 63
    1475 <212> TYPE: DNA
    1476 <213> ORGANISM: Artificial
W--> 1477 <220> FEATURE:
W--> 1478 <221> NAME/KEY:
    1479 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
    1480   RA patients
W--> 1481 <400> SEQUENCE: 118
    1483 tacttctgtg ccagcagttt ctgggcccct acggacaata atgagcagtt cttcgggcca   60
E--> 1484 gga   63 ← insert
    1529 <210> SEQ ID NO: 122
    1530 <211> LENGTH: 63
    1531 <212> TYPE: DNA
    1532 <213> ORGANISM: Artificial
W--> 1533 <220> FEATURE:
W--> 1534 <221> NAME/KEY:
    1535 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
    1536   RA patients
W--> 1537 <400> SEQUENCE: 122
    1539 tacttctgtg ccagcagccc tagggagggc ctcctcaata atgagcagtt cttcgggcca   60 .
E--> 1540 gga   63 ← insert
    1888 <210> SEQ ID NO: 148
    1889 <211> LENGTH: 61
    1890 <212> TYPE: DNA
    1891 <213> ORGANISM: Artificial
W--> 1892 <220> FEATURE:
W--> 1893 <221> NAME/KEY:
    1894 <223> OTHER INFORMATION: CDR3 nucleic acid sequence of BV14 clonotype derived from
ST specimen of
    1895   RA patients
W--> 1896 <400> SEQUENCE: 148
    1898 tacttctgtg ccagcagtcc ccgactagcg ggagataaag gagcagtact tcgggccggg   60 .
E--> 1899 c   61 ← insert

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Input Set : **A:\EP.txt**
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Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 21
Seq#:3; Line(s) 44
Seq#:4; Line(s) 58
Seq#:75; Line(s) 881
Seq#:77; Line(s) 910
Seq#:79; Line(s) 939
Seq#:81; Line(s) 967
Seq#:83; Line(s) 997
Seq#:85; Line(s) 1025
Seq#:87; Line(s) 1052
Seq#:89; Line(s) 1079
Seq#:91; Line(s) 1106
Seq#:93; Line(s) 1133
Seq#:95; Line(s) 1160
Seq#:97; Line(s) 1187
Seq#:99; Line(s) 1214
Seq#:101; Line(s) 1241
Seq#:103; Line(s) 1268
Seq#:105; Line(s) 1295
Seq#:107; Line(s) 1323
Seq#:109; Line(s) 1351
Seq#:111; Line(s) 1378
Seq#:113; Line(s) 1406
Seq#:115; Line(s) 1436
Seq#:117; Line(s) 1464
Seq#:119; Line(s) 1492
Seq#:121; Line(s) 1520
Seq#:123; Line(s) 1549
Seq#:125; Line(s) 1577
Seq#:127; Line(s) 1605
Seq#:129; Line(s) 1632
Seq#:131; Line(s) 1659
Seq#:133; Line(s) 1686
Seq#:135; Line(s) 1713
Seq#:137; Line(s) 1740
Seq#:139; Line(s) 1767
Seq#:141; Line(s) 1795
Seq#:143; Line(s) 1822
Seq#:145; Line(s) 1851
Seq#:147; Line(s) 1879
Seq#:149; Line(s) 1908
Seq#:151; Line(s) 1936
Seq#:153; Line(s) 1964

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Input Set : **A:\EP.txt**
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```
Seq#:155; Line(s) 1992
Seq#:157; Line(s) 2020
Seq#:159; Line(s) 2047
Seq#:161; Line(s) 2075
Seq#:163; Line(s) 2103
Seq#:165; Line(s) 2131
Seq#:167; Line(s) 2159
```

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

```
Seq#:1,2,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31
Seq#:32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,76,78,80,82,84
Seq#:86,88,90,92,94,96,98,100,102,104,106,108,110,112,114,116,118,120,122,124
Seq#:126,128,130,132,134,136,138,140,142,144,146,148,150,152,154,156,158,160
Seq#:162,164,166,168
```

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Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\07292003\J612468.raw**

```
L:7 M:283 W: Missing Blank Line separator, <120> field identifier
L:9 M:283 W: Missing Blank Line separator, <130> field identifier
L:10 M:283 W: Missing Blank Line separator, <140> field identifier
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:11 M:289 W: Identifier Missing or Out-Of-Order, <150> PRIOR APP NO
L:12 M:283 W: Missing Blank Line separator, <160> field identifier
L:18 M:283 W: Missing Blank Line separator, <220> field identifier
L:19 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:22 M:283 W: Missing Blank Line separator, <400> field identifier
L:30 M:283 W: Missing Blank Line separator, <220> field identifier
L:31 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:34 M:283 W: Missing Blank Line separator, <400> field identifier
L:42 M:283 W: Missing Blank Line separator, <220> field identifier
L:46 M:283 W: Missing Blank Line separator, <400> field identifier
L:56 M:283 W: Missing Blank Line separator, <220> field identifier
L:60 M:283 W: Missing Blank Line separator, <400> field identifier
L:69 M:283 W: Missing Blank Line separator, <220> field identifier
L:70 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:73 M:283 W: Missing Blank Line separator, <400> field identifier
L:82 M:283 W: Missing Blank Line separator, <220> field identifier
L:86 M:283 W: Missing Blank Line separator, <400> field identifier
L:109 M:283 W: Missing Blank Line separator, <220> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:137 M:283 W: Missing Blank Line separator, <220> field identifier
L:140 M:283 W: Missing Blank Line separator, <400> field identifier
L:148 M:283 W: Missing Blank Line separator, <220> field identifier
L:151 M:283 W: Missing Blank Line separator, <400> field identifier
L:159 M:283 W: Missing Blank Line separator, <220> field identifier
L:162 M:283 W: Missing Blank Line separator, <400> field identifier
L:170 M:283 W: Missing Blank Line separator, <220> field identifier
L:173 M:283 W: Missing Blank Line separator, <400> field identifier
L:181 M:283 W: Missing Blank Line separator, <220> field identifier
L:184 M:283 W: Missing Blank Line separator, <400> field identifier
L:192 M:283 W: Missing Blank Line separator, <220> field identifier
L:195 M:283 W: Missing Blank Line separator, <400> field identifier
L:203 M:283 W: Missing Blank Line separator, <220> field identifier
L:206 M:283 W: Missing Blank Line separator, <400> field identifier
L:215 M:283 W: Missing Blank Line separator, <220> field identifier
L:218 M:283 W: Missing Blank Line separator, <400> field identifier
L:226 M:283 W: Missing Blank Line separator, <220> field identifier
L:229 M:283 W: Missing Blank Line separator, <400> field identifier
L:237 M:283 W: Missing Blank Line separator, <220> field identifier
L:240 M:283 W: Missing Blank Line separator, <400> field identifier
L:248 M:283 W: Missing Blank Line separator, <220> field identifier
L:251 M:283 W: Missing Blank Line separator, <400> field identifier
L:259 M:283 W: Missing Blank Line separator, <220> field identifier
L:262 M:283 W: Missing Blank Line separator, <400> field identifier
L:270 M:283 W: Missing Blank Line separator, <220> field identifier
```

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Input Set : **A:\EP.txt**
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```
L:273  M:283 W: Missing Blank Line separator, <400> field identifier
L:281  M:283 W: Missing Blank Line separator, <220> field identifier
L:284  M:283 W: Missing Blank Line separator, <400> field identifier
L:292  M:283 W: Missing Blank Line separator, <220> field identifier
L:295  M:283 W: Missing Blank Line separator, <400> field identifier
L:303  M:283 W: Missing Blank Line separator, <220> field identifier
L:306  M:283 W: Missing Blank Line separator, <400> field identifier
L:784  M:252 E: No. of Seq. differs, <211> LENGTH:Input:21 Found:19 SEQ:66
L:896  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:76
L:925  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:78
L:953  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:80
L:982  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:82
L:1011 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:84
L:1039 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:86
L:1066 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:88
L:1093 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:90
L:1120 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:92
L:1147 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:94
L:1174 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:96
L:1201 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:98
L:1228 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:100
L:1254 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:102
L:1282 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:104
L:1310 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:106
L:1338 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:108
L:1365 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:110
L:1392 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:112
L:1421 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:114
L:1427 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:114
L:1450 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:116
L:1456 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:116
L:1478 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:118
L:1484 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:118
L:1507 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:120
L:1534 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:122
L:1540 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:63 SEQ:122
L:1563 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:124
L:1591 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:126
L:1619 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:128
L:1646 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:130
L:1673 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:132
L:1700 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:134
L:1727 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:136
L:1754 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:138
L:1781 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:140
L:1809 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:142
L:1837 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:144
L:1865 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:146
L:1893 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:148
```

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```
L:1899 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:61 SEQ:148
L:1923 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:150
L:1950 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:152
L:1978 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:154
L:2007 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:156
L:2034 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:158
L:2061 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:160
L:2089 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:162
L:2117 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:164
L:2145 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:166
L:2173 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:168
```